

SCORE Search Results Details for Application 09997683 and Search Result us-09-997-683-351.spdiraw.ra.

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This page gives you Search Results detail for the Application 09997683 and Search Result us-09-997-683-351.spdiraw.ra.

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GenCore version 5.1.9
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OM protein - protein search, using sw model

Run on: September 7, 2006, 00:45:10 ; Search time 50 Seconds
(without alignments)
344.871 Million cell updates/sec

Title: US-09-997-683-351
Perfect score: 1118
Sequence: 1 MPPAGLRRRAAPLTAIALLVL.....VYNPAAPPYPMPQPSYPGA 197

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 650591 seqs, 87530628 residues

Total number of hits satisfying chosen parameters: 650591

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1500 summaries

Database : Issued_Patents_AA:*
1: /EMC_Celerra_SIDS3/ptodata/2/iaa/5_COMB.pep:*
2: /EMC_Celerra_SIDS3/ptodata/2/iaa/6_COMB.pep:*
3: /EMC_Celerra_SIDS3/ptodata/2/iaa/7_COMB.pep:*
4: /EMC_Celerra_SIDS3/ptodata/2/iaa/H_COMB.pep:*
5: /EMC_Celerra_SIDS3/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /EMC_Celerra_SIDS3/ptodata/2/iaa/RE_COMB.pep:*
7: /EMC_Celerra_SIDS3/ptodata/2/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1118	100.0	197	2	US-09-991-181-351	Sequence 351, App
2	1118	100.0	197	2	US-09-990-444-351	Sequence 351, App
3	1118	100.0	197	2	US-09-997-333-351	Sequence 351, App
4	1118	100.0	197	2	US-09-992-598-351	Sequence 351, App
5	1118	100.0	197	2	US-09-989-735-351	Sequence 351, App
6	1118	100.0	197	3	US-09-989-726-351	Sequence 351, App
7	1118	100.0	197	3	US-09-997-514-351	Sequence 351, App
8	1118	100.0	197	3	US-09-989-728-351	Sequence 351, App
9	1118	100.0	197	3	US-09-997-349-351	Sequence 351, App
10	1118	100.0	197	3	US-09-997-653-351	Sequence 351, App